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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/582,277	06/10/2006	Allan Nielsen	10527.204-US 1622	
	7590 07/29/201 NORTH AMERICA,	EXAMINER		
500 FIFTH AVENUE SUITE 1600 NEW YORK, NY 10110			NOAKES, SUZANNE MARIE	
			ART UNIT	PAPER NUMBER
			1656	
			NOTIFICATION DATE	DELIVERY MODE
			07/29/2010	ELECTRONIC

Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.

Notice of the Office communication was sent electronically on above-indicated "Notification Date" to the following e-mail address(es):

Patents-US-NY@novozymes.com

	Application No.	Applicant(s)					
Office Action Comments	10/582,277	NIELSEN ET AL.					
Office Action Summary	Examiner	Art Unit					
	SUZANNE M. NOAKES	1656					
The MAILING DATE of this communication app Period for Reply	pears on the cover sheet with the c	orrespondence address					
A SHORTENED STATUTORY PERIOD FOR REPLY IS SET TO EXPIRE 3 MONTH(S) OR THIRTY (30) DAYS, WHICHEVER IS LONGER, FROM THE MAILING DATE OF THIS COMMUNICATION. - Extensions of time may be available under the provisions of 37 CFR 1.136(a). In no event, however, may a reply be timely filed after SIX (6) MONTHS from the mailing date of this communication. - If NO period for reply is specified above, the maximum statutory period will apply and will expire SIX (6) MONTHS from the mailing date of this communication. - Failure to reply within the set or extended period for reply will, by statute, cause the application to become ABANDONED (35 U.S.C. § 133). Any reply received by the Office later than three months after the mailing date of this communication, even if timely filed, may reduce any earned patent term adjustment. See 37 CFR 1.704(b).							
Status							
1)⊠ Responsive to communication(s) filed on <u>25 F</u> o	ehruary 2009						
<u>/_</u>	This action is FINAL . 2b) This action is non-final. Since this application is in condition for allowance except for formal matters, prosecution as to the merits is						
	closed in accordance with the practice under <i>Ex parte Quayle</i> , 1935 C.D. 11, 453 O.G. 213.						
closed in accordance with the practice under <i>Ex parte Quayre</i> , 1933 C.D. 11, 433 C.G. 213.							
Disposition of Claims							
4)⊠ Claim(s) <u>41 and 46-65</u> is/are pending in the ap	4)⊠ Claim(s) 41 and 46-65 is/are pending in the application.						
4a) Of the above claim(s) is/are withdraw	4a) Of the above claim(s) is/are withdrawn from consideration.						
5) Claim(s) is/are allowed.							
6)⊠ Claim(s) <u>41 and 46-65</u> is/are rejected.	·						
7) Claim(s) is/are objected to.							
8) Claim(s) are subject to restriction and/o	r election requirement.						
one conjust to realistic and conjust to realistic and real							
Application Papers							
9)☐ The specification is objected to by the Examiner.							
10) The drawing(s) filed on is/are: a) accepted or b) objected to by the Examiner.							
Applicant may not request that any objection to the drawing(s) be held in abeyance. See 37 CFR 1.85(a).							
Replacement drawing sheet(s) including the correction is required if the drawing(s) is objected to. See 37 CFR 1.121(d).							
11) The oath or declaration is objected to by the Examiner. Note the attached Office Action or form PTO-152.							
Priority under 35 U.S.C. § 119							
 12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f). a) All b) Some * c) None of: 1. Certified copies of the priority documents have been received. 2. Certified copies of the priority documents have been received in Application No. 3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)). * See the attached detailed Office action for a list of the certified copies not received. 							
Attachment(s) 1) Notice of References Cited (PTO-892) 2) Notice of Draftsperson's Patent Drawing Review (PTO-948) 3) Information Disclosure Statement(s) (PTO/SB/08) Paper No(s)/Mail Date	4) Interview Summary Paper No(s)/Mail Da 5) Notice of Informal P 6) Other:	te					

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DETAILED ACTION

1. Upon further consideration and in view of the Appeal Brief and filed on 19 May

2010, PROSECUTION IS HEREBY REOPENED. A new ground of rejection is set forth

below.

To avoid abandonment of the application, appellant must exercise one of the

following two options:

(1) file a reply under 37 CFR 1.111 (if this Office action is non-final) or a reply

under 37 CFR 1.113 (if this Office action is final); or,

(2) initiate a new appeal by filing a notice of appeal under 37 CFR 41.31 followed

by an appeal brief under 37 CFR 41.37. The previously paid notice of appeal fee and

appeal brief fee can be applied to the new appeal. If, however, the appeal fees set forth

in 37 CFR 41.20 have been increased since they were previously paid, then appellant

must pay the difference between the increased fees and the amount previously paid.

A Supervisory Patent Examiner (SPE) has approved of reopening prosecution by

signing below:

/Manjunath N. Rao /

Supervisory Patent Examiner, Art Unit 1656.

Status of the Claims

2. Claims 41 and 46-65 are pending and subject to examination on the merits.

Withdrawal of Previous Objections/Rejections

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3. As indicated in the Advisory action mailed 11/03/2009, the entry of Applicants amendments filed under 37 CFR 1.116 overcame all previous objections and 35 USC 112 1st paragraph rejections recited in the Final Office action (mail date 05/20/2009).

4. The rejection of claims 41 and 46-65 are rejected under 35 U.S.C. 102(b) as being anticipated by Chen et al. (Mol. Micro., 1995, cited on IDS) is withdrawn in favor of newly applied prior art. Appellants arguments in the Appeal Brief as filed 19 May 2010 are acknowledged but moot in view of the new rejections.

New Objections/Rejections

Claim Objections

5. Claims 47 and 59 are objected to under 37 CFR 1.75(c), as being of improper dependent form for failing to further limit the subject matter of a previous claim.

Applicant is required to cancel the claim(s), or amend the claim(s) to place the claim(s) in proper dependent form, or rewrite the claim(s) in independent form.

Independent claim 41 recites that the parent and progeny cells are both from the genus *Bacillus*. Thus, reiterating this in claim 47 is redundant.

Claim 59 recites that the progeny *Bacillus* cell is bacterial. This is inherent and redundant.

Claim Rejections - 35 USC § 102

6. The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

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A person shall be entitled to a patent unless -

(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

7. Claims 41, 46-49, 51, 52, 59-61, 63 and 64 are rejected under 35 U.S.C. 102(b) as being anticipated by Hartford and Dowds (Microbiology, 1994, Vol. 140, pp. 297-304) as evidenced by (Mol. Micro., 1995, cited on IDS).

Hartford and Dowds teach the isolation and characterization of hydrogen peroxide resistant mutant strains from *Bacillus subtilis*. The parent strain YB886 was exposed to increasing concentrations of hydrogen peroxide, several resistant cells survived and resulted in the creation of a new spontaneous progeny strain termed MA991. It is taught that several different proteins were over produced (termed overaccumulated in the reference) as compared to the wild-type parent strain. Specifically, Figure 5 shows the size of the proteins and N-terminal sequences of said proteins which were over-produced in the MA991 strain as compared to the parent strain YB886. It is noted that the 113 kDa protein, although it lacks the N-terminal methionine, is 100% identical to the amino acids 2-16 of the instant SEQ ID NO: 2. The next 10 amino acids, however, are different.

Hartford 113 kDa <u>? K T E N A K T N Q T L V E N</u> K S T T Q T V F R M H
SEQ ID NO: 2 M K T E N A K T N Q T L V E N S L N T Q L S N W F L

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Chen et al. teach using the MA991 strain as taught by Hartford and Dowds to make other progeny strains therefrom.

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It is specifically stated (see p. 297, 1st col., 1st paragraph, ½ way down to 2nd col., 1st paragraph):

"Hartford and Dowds (1994) isolated strain MA991 as a spontaneous H_2O_2 -resistant derivative of YB886, which constitutively synthesizes high levels of H_2O_2 -inducible proteins. These include catalase (KatA; 59.5 kDa) and alkyl hydroperoxide reductase (AhpC and AhpF; 23 and 53 kDa) (Fig. 4). In addition, MA991 overproduces two proteins with apparent molecular masses of 16 and 113 kDa and is reduced for flagellin (hag) expression.

We noticed that the published amino-terminal sequence of the 113 kDa protein overproduced in MA991 (Hartford and Dowds, 1994) is identical to MrgA for the first 15 of 26 amino acids. To test the relationship between *mrgA* and this 113 kDa protein, we transformed our *mrgA-lacZ* fusion into MA991 to generate strain HB1032. In stationary phase, HB1032 still overproduced KatA, AhpC and AhpF, but neither the 16 kDa nor the 113 kDa protein was observed by SDS-PAGE (Fig. 4). This suggests that *mrgA* is the structural gene for these two observed protein bands; the 113 kDa band presumably represents a stable oligomeric complex (see below) which the 16 kDa band is the appropriate size for the MrgA monomer. In support of this hypothesis, the aminoterminal sequence of the 16 kDa band was subsequently found to match exactly the predicted MrgA sequence (H. Cameron, unpublished data; cited in Dowds, 1994). This suggests that the late cycles of amino acid sequence of the 113 kDa protein were in error.

Previous pulse-labelling studies have identified a 16 kDa protein as the most strongly induced band following treatment of growing *B. subtilis* cells with 50 μM H₂O₂ (Murphy et al., 1987). Its synthesis is highest during the first 5 to 10 min after treatment with H₂O₂ and returns to the low basal level within 30 min (Dowds et al., 1987) consistent with the transient induction of *mrgA-lacZ* observed in our gene fusion experiments (Fig. 3). Therefore, we conclude that this 16 kDa band represents the MrgA monomer which then assembles into a stable, oligomeric complex detected as the 113 kDa band by SDS-PAGE."

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Therefore, Chen et al. make it sufficiently clear that the 113 kDa protein N-terminal sequence as taught by Hartford and Dowd originally was in error for amino acids 15-26 and evidences that said 113 kDa protein is inherently MrgA in oligomeric complex form. Said 113 kDa complex is clearly overproduced in MA991 as compared to YB886 as taught by Hartford and Dowd – see Figures 3 & 5, as are several other "proteins of interest" (e.g. KatA, AhpC and AhpF), .

It is further noted that step (b) as in claim 46, e.g. "recovering the protein" is met by the isolation of said protein on the noted SDS gels as well as the N-terminal sequencing of said protein complex.

8. Claims 41 and 46-65 are rejected under 35 U.S.C. 102(b) as being anticipated by Chen et al. (1995, PNAS, Vol. 140, pp. 297-304).

Chen et al. teach the creation of various strains of *Bacillus subtilis* which have been transformed with exogenous chromosomal mrgA-lacZ fusion DNA which is then fused to either the mrgA or katA promoters and expressed (see p. 8190 and p. 8192, as recited below). Said expression results in progeny cells which produce greater amounts of the fusion construct than the parent strain. It is specifically taught:

Each cured strain (HB13XX series) suspected of carrying a trans-acting mutation (because the constitutive phenotype was not phage-linked) was transduced to resistance to erythromycin and colincomycin by SP β 1122, generating strains HB12XXB, and constitutive expression of β -gal was confirmed. To study the regulation of mrgA and mrgC (1) in these mutant backgrounds, each HB13XX strain was transformed with HB1022(mrgA-lacZ)chromosomal DNA (1), generating strains HB14XX, or was transduced with SP β 085(mrgC-cat-lacZ), generating strains HB15XX. – See p. 8190, 2nd col., last paragraph.

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And (see p. 8192, 1st col., 2nd and 3rd paragraphs):

Characterization of Trans-Acting Mutations. To define factors involved in the regulation of mrgA, we have characterized 12 trans-acting mutants. Each mutant was cured of the SP β 1122 prophage, and transcriptional fusions to the mrgA (HB14XX series) and mrgC (HB15XX series) promoters were introduced. All 12 HB14XX strains displayed increased mrgA-lacZ expression, which was repressed little, if at all, by addition of Mn(II) (data not shown). We have described a second metalloregulated gene in *B. subtilis*, mrgC, which is repressed by iron but not by Mn(II) (1). In all 12 HB15XX strains, both mrgC-lacZ expression and the synthesis of catecholate siderophores were repressed normally by iron (data not shown). Therefore, regulation of mrgA expression [by both Mn(II) and iron] is independent of the postulated iron-dependent repressor, which regulates siderophore biosynthesis and mrgC expression.

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Like our trans-acting mutants, an H202R strain isolated previously, MA991 (8), is derepressed for mrgA expression. MA991 has a characteristic protein profile when analyzed by Coomassie-stained SDS/PAGE: MrgA, KatA, AhpC, and AhpF are all overproduced (8, 42). Nine of our trans-acting mutants shared this altered protein profile (data not shown) and are therefore likely to be mutant in the same regulatory pathway or even the same gene.

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Therefore, the protein of interest can be the β-galactosidase although it is apparent that said protein of interest can also be KatA, AhpC or AhpF as it is taught that these proteins are all over expressed as well. (see above and also Figure 4, Group I strains).

Reference of Interest – Not Relied Upon

9. Perkins et al. "Construction and properties of Tn917-lac, a transposon derivative that mediates transcriptional gene fusions in *Bacillus subtilis*", PNAS, 1986, Vol 83, pp. 140-144 – describes the TN917-lacZ fusion construct which is capable of generating fusions that connect the transcripts of *Bacillus subtilis* chromosomal genes to the coding sequence of the lacZ gene of *Escherchia coli*.

Conclusion

- 10. No claim is allowed.
- 11. Any inquiry concerning this communication or earlier communications from the examiner should be directed to SUZANNE M. NOAKES whose telephone number is (571)272-2924. The examiner can normally be reached on 7.00 AM-3.30 PM.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Manjunath Rao can be reached on 571-272-0939. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

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Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free). If you would like assistance from a USPTO Customer Service Representative or access to the automated information system, call 800-786-9199 (IN USA OR CANADA) or 571-272-1000.

/SUZANNE M. NOAKES/ Primary Examiner, Art Unit 1656 20 July 2010